

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

## Histone Database

RRID:SCR\_007711

Type: Tool

### Proper Citation

Histone Database (RRID:SCR\_007711)

### Resource Information

**URL:** <http://research.nhgri.nih.gov/histones/>

**Proper Citation:** Histone Database (RRID:SCR\_007711)

**Description:** Histone Database is a database of histones and their corresponding sequences. Sequence- and text-based searches were performed on NCBI's redundant and non-redundant (nr) peptide sequence databases. These databases are derived from GenBank, EMBL, and DDBJ translated DNA coding regions, plus protein sequences from the PDB (Protein Data Bank), SWISS-PROT, the PIR (Protein Information Resource), and the PRF (Protein Research Foundation). :Users can search by keyword, sequence fragment, category, organism, and redundancy of the set.

**Synonyms:** Histone Database

**Resource Type:** data or information resource, database

**Keywords:** histone

**Resource Name:** Histone Database

**Resource ID:** SCR\_007711

**Alternate IDs:** nif-0000-02961

**Record Creation Time:** 20220129T080243+0000

**Record Last Update:** 20260729T044154+0000

### Ratings and Alerts

No rating or validation information has been found for Histone Database.

No alerts have been found for Histone Database.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 5 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [nidm-terms](#) .

Anderson L, et al. (2012) Schistosoma mansoni histones: from transcription to chromatin regulation; an in silico analysis. Molecular and biochemical parasitology, 183(2), 105.

Mariño-Ramírez L, et al. (2011) The Histone Database: an integrated resource for histones and histone fold-containing proteins. Database : the journal of biological databases and curation, 2011, bar048.

Malik A, et al. (2007) Databases and QSAR for cancer research. Cancer informatics, 2, 99.

Gissot M, et al. (2007) Epigenomic modifications predict active promoters and gene structure in Toxoplasma gondii. PLoS pathogens, 3(6), e77.

Galperin MY, et al. (2005) The Molecular Biology Database Collection: 2005 update. Nucleic acids research, 33(Database issue), D5.